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<!--StartFragment-->US-10-179-373-17
; Sequence 17, Application US/10179373
; Publication No. US20030232407A1
; GENERAL INFORMATION:
; APPLICANT: ZOLLER, MARK
; APPLICANT: LI, XIAODONG
; APPLICANT: STASZEWSKI, LENA
; APPLICANT: O'CONNELL, SHAWN
; APPLICANT: ZOZULYA, SERGEY
; APPLICANT: ADLER, JON
; APPLICANT: XU, HONG
; APPLICANT: ECHEVERRI, FERNANDO
; TITLE OF INVENTION: T1R HETERO-OLIGOMERIC TASTE RECEPTORS AND CELL LINES
; TITLE OF INVENTION: THAT EXPRESS SAID RECEPTORS AND USE THEREOF FOR
; TITLE OF INVENTION: IDENTIFICATION OF TASTE COMPOUNDS
; FILE REFERENCE: 078003-0291566
; CURRENT APPLICATION NUMBER: US/10/179,373
; CURRENT FILING DATE: 2002-06-26
; PRIOR APPLICATION NUMBER: 60/300,434
; PRIOR FILING DATE: 2001-06-26
; PRIOR APPLICATION NUMBER: 60/304,749
; PRIOR FILING DATE: 2001-07-13
; PRIOR APPLICATION NUMBER: 60/310,493
; PRIOR FILING DATE: 2001-08-08
; PRIOR APPLICATION NUMBER: 60/331,771
; PRIOR FILING DATE: 2001-11-21
; PRIOR APPLICATION NUMBER: 60/339,472
; PRIOR FILING DATE: 2001-12-14
; PRIOR APPLICATION NUMBER: 60/372,090
; PRIOR FILING DATE: 2002-04-15
; PRIOR APPLICATION NUMBER: 60/374,143
; PRIOR FILING DATE: 2002-04-22
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 17
; LENGTH: 843
; TYPE: PRT
; ORGANISM: Rattus sp.
US-10-179-373-17
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Query Match          91.9%; Score 4134; DB 4; Length 843;
Best Local Similarity 91.0%; Pred. No. 0;
Matches 767; Conservative 34; Mismatches 42; Indels 0; Gaps 0;
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Db      1 MGQPQARTLCLLSLLHVLKPKGKLVENSDFHLAGDYLLGGLFTLHANVKSISHLSYLQVP 60

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Db     61 KCNEFTMKVLGYNLMQAMRFAVEEINNCSLLPGVLLGYEMVDVCYLSNNIHPGLYFLAQ 120

Qy    121 IDDFLPILKDYSQYRPQVAVIGPDNSESATVSNILSYFLVPQVTYSAITDKLQDKRRF 180
Db    121 DDDLPLILKDYSQYMPHVAVIGPDNSESATVSNILSHFLIPQITYSAISDKLRDKRRF 180

Qy    181 PAMLRTPVSATHHIEAMVQLMVHFPQWNWIVVLVSDDDYGRENSHLLSQRLTNTGDICIAF 240
Db    181 PSMLRTPVSATHHIEAMVQLMVHFPQWNWIVVLVSDDDYGRENSHLLSQRLTKTSDICIAF 240

Qy    241 QEVLVPPEPNQAVRPEEQQLDNILDKLRRTSARVVVIFSPELSLHNFFREVLRWNFTGF 300
Db    241 QEVLPIPESSQVMRSEEQRLDNILDKLRRTSARVVVIFSPELSLYSPFHEVLRWNFTGF 300

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Qy    361 TTNQDCDACMNNITESFNNVLMLSGERVVYSVYSAVAVAHTLHRLHLCNQVRCTKQIVY 420
Db    361 TTNQDCDACLNTTKSFNNILILSGERVVYSVYSAVAVAHALHRLGCVNRVCTKQKVY 420

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Db    421 PWQLLREIWHVNFTLLGNRLFFDQGDMPMLLDIIQWQWGLSQNPFSIASYSPTSKRLT 480

Qy    481 YISNVSWYTPNNTVPISMCSKSCQPGQMKKPIGLHPCCFECDPCPDYLNRSVDEFNCL 540
Db    481 YINNVSWYTPNNTVPSMCSKSCQPGQMKKSVGLHPCCFECLDCMPGYLNRSADFNCL 540

Qy    541 SCPGSMWSYKNNIACPKRRLAFLEWHEVPTIVVTILAALGFISTLAILLIFWRHFQTPMV 600
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Db      601 RSAGGPMCFLMLVPLLLAFGMVPVYVGPPTVFSCFCRQAFFTVCFSICLSCITVRSPQIV 660
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Db      661 CVFKMARRLPSAYSFWWMRYHGPYVVFVAFITAIKVALVVGNNMLATTINPIGRTPDDPNIM 720
Qy      721 ILSCHPNYRNGLLFNNTSMDLLSVLGFSFAYVGKELPTNYNEAKFITLSMTFSPTSSISL 780
Db      721 ILSCHPNYRNGLLFNNTSMDLLSVLGFSFAYMGKELPTNYNEAKFITLSMTFSPTSSISL 780
Qy      781 CTFMSVHDGVLVTIMDLLVTVLNFLAIGLGYFGPKCYMILFYPERNTSAYFNMSMIQGYTM 840
Db      781 CTFMSVHDGVLVTIMDLLVTVLNFLAIGLGYFGPKCYMILFYPERNTSAYFNMSMIQGYTM 840
Qy      841 RKS 843
Db      841 RKS 843
    
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; SEQ ID NO 17
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US-10-179-373-17
    
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Query Match 100.0%; Score 4494; DB 4; Length 843;
 Best Local Similarity 100.0%; Pred. No. 0;
 Matches 843; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db	1	MGPQARTLCLLSLLHLVLPKPGKLVENSDFHLAGDYLLGGLFTLHANVKSISHLSYLVQVP	60
Qy	61	KCNEFTMKVLGYNLMQAMRFAVEEINNCSSLLPGVLLGYEMVDVCYLSNNIHPGLYFLAQ	120
Db	61	KCNEFTMKVLGYNLMQAMRFAVEEINNCSSLLPGVLLGYEMVDVCYLSNNIHPGLYFLAQ	120
Qy	121	DDDLLPILKDYSQYMPHVAVIGPDNSESATVSNILSHFLIPQITYSAISDKLRDKRHF	180
Db	121	DDDLLPILKDYSQYMPHVAVIGPDNSESATVSNILSHFLIPQITYSAISDKLRDKRHF	180
Qy	181	PSMLRTVPSATHHIEAMVQLMVHFQWNWIVLVSDDDYGRENSHLLSQRLTKTSDICIAF	240
Db	181	PSMLRTVPSATHHIEAMVQLMVHFQWNWIVLVSDDDYGRENSHLLSQRLTKTSDICIAF	240
Qy	241	QEVLPPIESSQVMRSEEQRLDNILDKLRRTSARVVVVFSPELSLYSFPHEVLRWNFTGF	300
Db	241	QEVLPPIESSQVMRSEEQRLDNILDKLRRTSARVVVVFSPELSLYSFPHEVLRWNFTGF	300
Qy	301	VWIASSEWAIDPVLHNLTELRTGTFLGVTIQVRSIPGFSQFRVRRDKPGYPVPNTTNLR	360
Db	301	VWIASSEWAIDPVLHNLTELRTGTFLGVTIQVRSIPGFSQFRVRRDKPGYPVPNTTNLR	360
Qy	361	TTCNQDCDACLTNTKSFNNILILSGERVVYSVYSAVYAVAHALHRLGCMNRVCTKQKVY	420
Db	361	TTCNQDCDACLTNTKSFNNILILSGERVVYSVYSAVYAVAHALHRLGCMNRVCTKQKVY	420
Qy	421	PWQLLREIWHVNFTLLGNRLFFDQQGDMPLLDIIQWQWDLSONPFFQSIASYSPTSKRLT	480
Db	421	PWQLLREIWHVNFTLLGNRLFFDQQGDMPLLDIIQWQWDLSONPFFQSIASYSPTSKRLT	480
Qy	481	YINNVSWYTPNNTVPVSMCSKSCQPGQMKS SVGLHPCCFECLDCMPGTYLNRSADEFNCL	540
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Qy      601 RSAGGPMCFLMLVPLLLAFGMVPVYVGPPTVFSCFCRQAFPTVCFSICLSCITVRSPQIV 660
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Qy      661 CVFKMARRLPSAYSFWMRYHGPYVFVAFITA IKVALVVG NMLATTINPIGRTPDDPNIM 720
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Qy      721 ILSCHPNYRNGLLFN TSM D L L S V L G F S F A Y M G K E L P T N Y N E A K F I T L S M T F S F T S S I S L 780
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Qy      781 CTFMSVHDGVLVTIMDLLVTVLNFLAIGLGYFGPKCYMILFYPERNTSAYFN SMIQGYTM 840
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Qy      841 RKS 843
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Db      841 RKS 843
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